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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Perfect score:
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                      1008
3900
3900
363.5
34.5
218.3
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21.5
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1: /cgn2 6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

2: /cgn2 6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

3: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

4: /cgn2 6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

5: /cgn2 6/ptodata/2/pubpaa/US07 NEW PUB.pep:*

6: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

7: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

9: /cgn2 6/ptodata/2/pubpaa/US08 NEW PUB.pep:*

9: /cgn2 6/ptodata/2/pubpaa/US09A PUBCOMB.pep:*

10: /cgn2 6/ptodata/2/pubpaa/US09A PUBCOMB.pep:*

11: /cgn2 6/ptodata/2/pubpaa/US09A PUBCOMB.pep:*

12: /cgn2 6/ptodata/2/pubpaa/US09A PUBCOMB.pep:*

13: /cgn2 6/ptodata/2/pubpaa/US09 NEW PUB.pep:*

14: /cgn2 6/ptodata/2/pubpaa/US10A PUBCOMB.pep:*

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17: /cgn2 6/ptodata/2/pubpaa/US10C PUBCOMB.pep:*

18: /cgn2 6/ptodata/2/pubpaa/US10C PUBCOMB.pep:*

16: /cgn2 6/ptodata/2/pubpaa/US10C PUBCOMB.pep:*

17: /cgn2 6/ptodata/2/pubpaa/US10C PUBCOMB.pep:*

18: /cgn2 6/ptodata/2/pubpaa/US10C PUBCOMB.pep:*
                                                                                                                                                                                                                                                                                        Query
Match
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                                         US-09-966-880A-8
1086
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US-10-460-923-4
US-10-460-923-6
US-10-460-923-6
US-10-460-923-6
US-10-460-923-6
US-10-157-031-14
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US-10-166-880A-36
US-10-104-047-3729
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US-10-460-923-5
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Sequence 2, Appli
Sequence 5, Appli
Sequence 174, App
Sequence 174, Appli
Sequence 1639, Appli
Sequence 7, Appli
Sequence 159, Appli
Sequence 159, Appli
Sequence 16, Appli
Sequence 14, Appli
Sequence 14, Appli
Sequence 14, Appli
Sequence 14, Appli
Sequence 3, Appli
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	68346	48829,	42	122, Ap	61, App	65, App	55	615, Ap	63, App	Sequence 63, Appl	4, Appl	25, App	47947,	U L	143617,	164, Ap	135, Āp	e 14, App	6748	955, App	s 5, Appl	e 5, Appl	e 5, Appl	15, Appl	nce 3, App	equence 3, Appl	ce 3, Appl	quence 38853, A	quence 34155,	equence 77, App	

ALIGNMENTS

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CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: PCT/JP00/01918
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: JP 11-371382
PRIOR FILING DATE: 1999-12-27
PRIOR PPLICATION NUMBER: JP 11-178999
PRIOR FILING DATE: 1999-06-24
PRIOR APPLICATION NUMBER: JP 11-87192
PRIOR APPLICATION NUMBER: JP 11-87192
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 198
TYPE: PAT
ORGANISM: Homo sapiens
US-09-966-880A-8
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                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: HONJO, TASUKU
APPLICANT: MUZAMATSU, MASAMICHI
TITLE OF INVENTION: NOVEL CYTIDINE DEAMINASE
FILE REFERENCE: 06501-088001
CURRENT APPLICATION NUMBER: US/09/966,880A
CURRENT FILING DATE: 2001-09-28
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61
                                FLRYISDWDLDPGRCYRVTWFTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYFCEDRK 120
                                                                                                          MDSLLMNRKFLYQFKNVRWAKGRRETYLCYVVKRRDSATSFSLDFGYLRNKNGCHVELL
                                                                                MDSLLMNRRKFLYQFKNVRWAKGRRETYLCYVVKRRDSATSFSLDFGYLRNKNGCHVELL
                                                                                                                                                                   100.0%; Score 1086; DB 9; ilarity 100.0%; Pred. No. 8.1e-114; Conservative 0; Mismatches 0;
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PRIOR APPLICATION NUMBER: PCT/JP00/01918
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: JP 11-371382
PRIOR FILING DATE: 1999-12-27
PRIOR FILING DATE: 1999-12-27
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: JP 11-87192
PRIOR APPLICATION NUMBER: JP 11-87192
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
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Best Local Similarity
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    APPLICANT: MALIM, Michael H.
APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S.
APPLICANT: HARRIS, Reuben S.
APPLICANT: BISHOP, Kate N.
APPLICANT: NEUBERGER, Michael S.
APPLICANT: GADDIS, Nathan C.
APPLICANT: GADDIS, Nathan C.
APPLICANT: SIXON, James H.M.
TITLE OF INVENTION: DNA Deamination Mediates Innate Immunity to Retroviral Infection
FILE REFERENCE: 2225-74380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Honjo, Tasuku
APPLICANT: Muzamatsu, Masamichi
TITLE OF INVENTION: NOVEL CYTIDINE DEAMINASE
FILE REFERENCE: 06501-088001
CURRENT APPLICATION NUMBER: US/09/966,880A
CURRENT FILING DATE: 2001-09-28
CURRENT APPLICATION NUMBER: US/10/460,923
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ORGANISM: Mus musculus
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Pred. No. 4.8e-105;
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; ORGANISM: Homo sapiens US-09-729-674-174
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; ORGANISM: Homo sapiens
US-10-460-923-5
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Patent No. US20010039335A1
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                                                                  Query Match 35.9%;
Best Local Similarity 44.9%;
Matches 83; Conservative 3
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Best Local (
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PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 60/472,952
PRIOR FILING DATE: 2003-05-23
NUMBER OF SEQ ID NOS: 12
SOFTMARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                         APPLICANT: Fechtel, Kim

APPLICANT: Genetics Institute, Inc.
APPLICANT: Genetics Institute, Inc.
TITLE OF INVENTION: SECRETED PROTEINS AND POLYNUCLEOTIDES ENCODING THEM
FILE REFERENCE: 6055-64X
CURRENT APPLICATION NUMBER: US/09/729,674
CURRENT FILING DATE: 2000-12-04
PRIOR APPLICATION NUMBER: 09/539,330
PRIOR FILING DATE: 2000-03-30
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
EQ ID NO 174
LENGTH: 384
                                                                                                                                                                                                                                                                                        PRIOR FILING DATE: 2000-03-30 NUMBER OF SEQ ID NOS: 283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Jacobs,
                                                                                                                                                                                                   LENGTH: 384
TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          121 -DÓQGRCQÉGLRTLAEÁGAKÍSIMTYSEFKHŐMDTFVDHQGCPFQPWDGLDEHSQDLSGR 179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 62 HABICFLDVÍPFWKLDLDQDÝRVÍCFTSWSFCFSCAQEMAKFISKNKHVSLCÍFTARIÝ- 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ch 35.9%; Score 390; DB 15;
l Similarity 44.9%; Pred. No. 1.5e-35;
83; Conservative 31; Mismatches 59;
6 MNRRKFLYQPKAVRWAKGRRETYLCYVVKRRDSATSFSLD--FGYLRNK------NGC 55
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Steininger II, Rober
Spaulding, Vikki
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LaVallie, Edward R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Merberg, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Evans, Cheryl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wong,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gordon G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hilary
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Robert J.
                                                                  %; Score 390; DB 9; Length 384
%; Pred. No. 3.8e-35;
31; Mismatches 59; Indels
                                                                                                            Length 384;
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US-09-925-300-1639
, Sequence 1639, Application US/09925300
; Patent No. US20020151681A1
; GENERAL INFORMATION:
APPLICANT: Craig Rosen,
APPLICANT: Steve Ruben
; TITLE OF INVENTION: Nucleic Acids, Protein Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR FILING DATE: 2002-05-23
PRIOR FILING DATE: 2003-05-23
NUMBER OF SEQ ID NOS: 12
SOPTWARE: Patentin version 3.1
SEQ ID NO 2
LENGTH: 384
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US-10-460-923-2
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: GADDIS, Nathan C.
APPLICANT: SIMON, James H.M.

TITLE OF INVENTION: DNA Deamination Mediates Innate Immunity to Retroviral Infection
FILE REFERENCE: 22253-74380

CURRENT APPLICATION NUMBER: US/10/460,923

CURRENT FILING DATE: 2003-06-13

CURRENT FILING DATE: 2003-06-13
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APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S.
APPLICANT: BISHOP, Kate N.
APPLICANT: BUSBERGER, Michael S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            83;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               LRRIL 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -DDQGRCQEGLRTLAEAGAKISIMTYSEFKHCWDTFVDHQGCPFQPWDGLDEHSQDLSGR 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CEDRKAEPEGLRRLHRAGVQIAIMTFKDYFYCWNTFVENHERTFKAWEGLHENSVRLSRQ 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HAELCFLDVIPFWKLDLDQDYRVTCFTSWSPCFSCAQEMAKFISKNKHVSLCIFTARIY- 315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HVELLFLRYISDWDLDPGRCYRVTWFTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYF 115
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                                                                                                                                                                                                                                                                                                                                                                                   LRAIL 379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Score 390; DB 15; ; Pred. No. 3.8e-35; 31; Mismatches 59;
                                             Proteins and Antibodies
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CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05988
PRIOR FILING DATE: 2000-03-08
PRIOR PLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1890
SEQ ID NO 1639
LENGTH: 222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S.
APPLICANT: HIGHOP, Kate N.
APPLICANT: BISHOP, Kate N.
APPLICANT: GADDIS, Mathan C.
APPLICANT: GADDIS, Nathan C.
APPLICANT: SIMON, James H. M.
TITLE OF INVENTION: DNA Deamination Mediates Innate Immunity to Retroviral Infection FILE REFERENCE: 22253-74380
CURRENT APPLICATION NUMBER: US/10/460,923
CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR APPLICATION NUMBER: US 60/472,952
RNOR FILING DATE: 2003-06-13
PRIOR FILING DATE: 2003-05-23
NUMBER OF SEQ ID NOS: 12
SOOTWARE: Patentin version 3.1
SEQ ID NO 7
1FROTH. 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/10460923
Publication No. US20040009951A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: MALIM, Michael H.
APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S
APPLICANT: BISHOP, Kate N.
APPLICANT: NEUBERGER, Michae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH: 199
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   113 LYFCEDRKAEP---EGIRRIHRAGVQIAIMTFKDYFYCWNTFVENHERTFKAWEGIHENS 169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                125 GLRRLHRAGVQIAIMTFKDYFYCWNTFVENHERTFKAWEGLHENSVRLSRQLRRIL 180
                                                                                                             69 RHABLRFLDLVPSLQLDPAQIYRVTWFISWSPCFSWGCAGEVRAFLQENTHVRLRIFAAR
                                                                                                                                                               55 CHVELLFLRYISDWDLDPGRCYRVTWFTSWSPCYD--CARHVADFLRGNPNLSLRIFTAR 112
                                                                                                                                                                                                                                            12 LMDPHIFTSNFNN---GIGRHKTYLCYEVERLDNGTSVKMDQHRGFLHNQAKNLLCGFYG
                                                                                                                                                                                                                                                                                                                                                                            83;
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                                                                                                                                                                                                                                                                                                        5 LMNRRKFLYQFKNVRWAKGRRETYLCYVVKRRDSATSFSLD--FGYLRN--KN-----
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83; Conservative
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Similarity 44.9%;
79; Conservative 2:
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                                                                                                                                                                                                                                                                                                                                                                     Score 363.5; DB 15;
Pred. No. 1.6e-32;
B; Mismatches 57;
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Pred. No. 3.8e-33;
4; Mismatches 64
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CURRENT APPLICATION NUMBER: US/10/460,923
CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 60/472,952
PRIOR FILING DATE: 2003-05-23
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                 US-10-247-671-159
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TYPE: PRT
ORGANISM: Unknown
FEATURE:
OTHER INFORMATION: mouse orthologue
US-10-460-923-4
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APPLICANT: SHEEH
APPLICANT: HARRI
APPLICANT: BISHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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APPLICANT: Shiffman, Dov
APPLICANT: Porter, Gordon, J.
APPLICANT: Waser, Matthew R.
TITLE OF INVENTION: GENES EXPRESSED IN TREATED FOAM CELLS
FILE REFERENCE: PA-0050 US
CURRENT APPLICATION NUMBER: US/10/247,671
CURRENT FILING DATE: 2002-09-18
PRIOR APPLICATION NUMBER: 60/323,784
PRIOR FILING DATE: 2001-09-19
                                                                                                                                                          APPLICANT: Mikit
APPLICANT: Shif
APPLICANT: Port
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: NEUBERGER, Michael S.
APPLICANT: GADDIS, Nathan C.
APPLICANT: SIMON, James H.M.
TITLE OF INVENTION: DNA Deamination Mediates Innate Immunity to Retroviral Infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 22253-74380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity es 71; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         80 FHDKVLKVLSPREEFKITWYMSWSPCFECAEQVLRFLATHHNLSLDIFSSRLYNIRDPEN 139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EPEGIRRLHRAGVQIAIMTFKDYFYCWNTFVENHERTFKAWEGLHENSVRLSRQLRRILL 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ISD---WDLDPGRCYRVTWFTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYFCEDRKA 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LISQETFKFHFKNLRYAIDRKDTFLCYEVTRKDCDSPVSLHHGVFKNKDNIHAEICFLYW 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QALSGRIRAIL 194
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HARRIS, Reuben S.
BISHOP, Kate N.
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                                                                                                                                                                                                                                                   Application US/10247671
o. US20030194721A1
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                                                                                                                                                                                                          Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         32.1%; Score 349; DB 15; 38.8%; Pred. No. 7.2e-31; tive 36; Mismatches 72;
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; PEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20030194721A1 135626CD1
US-10-247-671-159
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                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: PRT; ORGANISM: Homo sapiens US-10-460-923-3
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SOFTWARE: PERL Program
SEQ ID NO 159
LENGTH: 152
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.1 SEQ ID NO 3
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Tocal Similarity
                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: GADDIS, Nathan C.
APPLICANT: SIMON, James H.M.
TITLE OF INVENTION: DNA Deamination Mediates Innate Immunity to Retroviral Infection FILE REFERENCE: 22253-74380
CURRENT APPLICATION NUMBER: US/10/460,923
CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR FILING DATE: 2002-05-23
PRIOR FILING DATE: 2003-05-23
PRIOR SECTION NUMBER: US 60/472,952
PRIOR FILING DATE: 2003-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MALIM, Michael H.
APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S.
APPLICANT: BISHOP, Kate N.
APPLICANT: NEUBERGER, Michae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 195
                                                                                                                                                                                                                                                                                                                                               Local
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                                                                                              121 AEPEGIRRI--HRAG--VQIAIMTFKDYFYCWNTFVENHERTFKAWEGIHENSVRISRQI 176
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                                                                                                                                                                                  62 LRYISDW-DLDPGRCYRVTWFTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYFCEDRK 120
                                                                                                                                                                                                                                12 MYRDTFSYNFYNRPILSRRNTVWLCYEVKTKGPSRP-PLDAKIFRGQVYSELKYHPEMRF 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97
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                                                                                                                                            71 FHWFSKWRKLHRDQEYEVTWYISWSPCTKCTRDMATFLAEDPKVTLTIFVARLYYFWDPD
                                                                                                                                                                                                                                                                            6 MNRRKFLYQFKNVRWAKGRRETYLCYVVKRRDSATSFSLDF----GYLRNKNGCHVELLF
                                                                                                                                                                                                                                                                                                                            63;
                                                                                                                                                                                                                                                                                                                       h 22.0%; Score 238.5; DB 15;
Similarity 34.2%; Pred. No. 1.8e-18;
63; Conservative 26; Mismatches 84;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SHEEHY, Ann M.
HARRIS, Reuben S.
BISHOP, Kate N.
NEUBERGER, Michael S.
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                                                  YQ-EALRSICOKRDGPRATMKIMNYDEFOHCWSKFVYSQRELFEPWNNLPKYYILLHIMI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             43;
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                                                                                                                                                                                                                                                                                                                                                                     Length 195;
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Gaps

9

189

130

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Sequence 14, Application US/10157031

Publication No. US20030108890A1

GENERAL INFORMATION:
APPLICANT: Baranova, A. V.
APPLICANT: Yankovsky, N. K.
APPLICANT: Kozlov, A. P.
APPLICANT: Kozlov, A. P.
APPLICANT: Kozlov, A. P.
APPLICANT: Krukovskaya, L. L.
TITLE OF INVENTION: In silico screening for phenotype-associated explication application NUMBER: US/10/157,031

CURRENT APPLICATION NUMBER: US/10/157,031

CURRENT FILING DATE: 2002-05-30

NUMBER OF SEG ID NOS: 415

SOFTWARE: Patentin version 3.1
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US-10-157-031-14
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SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 6
LENGTH: 219
TYPE: PRT
ORGANISM: Unknown
FEATURE:
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US-10-460-923-6
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CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR APPLICATION NUMBER: US 60/472,952
PRIOR APPLICATION NUMBER: US 60/472,952
PRIOR FILING DATE: 2003-05-23
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CURRENT APPLICATION NUMBER: 110/10/100
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APPLICANT: SHEEHY, Ann M.
APPLICANT: HARRIS, Reuben S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 33.5%;
hes 67; Conservative --
                                                                                                                                                                                                                                                                                                                                                                                                                 196 İ-KESWGLQDLVNDFGNLQL 214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              138 RPFO-KGLCSLWOSGILVDVMDLPQFTDCWTNFV-NPKRPFWPWKGLEIISRRTQRRLHR 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                119 RKAEPEGLRRLHRAGVQIAIMTFKDYFYCWNTFVENHERTFKAWEGLHENSVRLSRQLRR 178
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HARRIS, Reuben S.
BISHOP, Kate N.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 230; DB 15;
Pred. No. 1.9e-17;
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; ORGANISM: Homo sapiens
US-10-157-031-14
RESULT 14
US-09-966-880A-36
                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 236
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-10-460-923-8
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US-10-460-923-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 8
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Best Local :
                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                      Query Match
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APPLICANT: SIMON, James H.M.
TITLE OF INVENTION: DAN Deamination Mediates Innate Immunity to Retroviral Infection
FILE REFERENCE: 22253-74380
CURRENT APPLICATION NUMBER: US/10/460,923
CURRENT FILING DATE: 2003-06-13
PRIOR APPLICATION NUMBER: US 60/388,513
PRIOR FILING DATE: 2002-06-13
PRIOR FILING DATE: 2002-06-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/472,952
PRIOR FILING DATE: 2003-05-23
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin version 3.1
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                                                                                                                                                                                                                                                                                                                        Local
                                                                          152
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                                                                                                                                                   93 CWECSQAIREFLSRHPGVTLVIYVARLFWHMDQQ-NRQGLRDLVNSGVTIQIMRASEYYH
                                                                                                                                                                                       87 CYDCARHVADFLRGNPNLSLRIFTARLYFCEDRKAEPEGLRRLHRAGVQIAIMTFKDYFY 146
                                                                                                                                                                                                                           33 RKEACLLYEIKWGMSRKIWRSSGKNTTNHVEVNFIKKFTSERDFHPSISCSITWFLSWSP
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                                                                                                              CWNTFV 152
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                                                                          CWRNEV 157
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SHEEHY, Ann M.
HARRIS, Reuben S.
BISHOP, Kate N.
NEUBERGER, Michael S.
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                                                                                                                                                                                                                                                                                                                      19.9%; Score 216.5; DB 15; 34.9%; Pred. No. 6.8e-16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20.1%; Score 218.5; DB 34.9%; Pred. No. 4e-16;
                                                                                                                                                                                                                                                                                                    31;
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                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                        Length 236;
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; Sequence 36, Application ; Patent No. US20020164743A ; GENERAL INFORMATION:

US/09966880A

APPLICANT: Honjo, Tasul APPLICANT: Muramatsu,

Tasuku 164743A1

Masamichi

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Sequence 3729, Application US/10104047

Sequence 3729, Application US/2001036392A1

GENERAL INFORMATION: US20030236392A1

APPLICANT: HELIX RESEARCH INSTITUTE
ITILE OF INVENTION: No. US20030236392A1e1 full length cDNA
FILE REFERENCE: H1-A0105

CURRENT APPLICATION NUMBER: US/10/104,047

CURRENT FILING DATE: 2002-03-25

PRIOR FILING DATE: PRIOR APPLICATION NUMBER:

PRIOR FILING DATE:

NUMBER OF SEQ ID NO 3729

SEQ ID NO 3729

LENGTH: 127

TYPE: DETTION DETTION OF SEQ ID NO 3729

LENGTH: 127
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US-10-104-047-3729
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FILE REFERENCE: 06501-088001
CURRENT APPLICATION UNMBER: US/09/966,880A
CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION UNMBER: PCT/JD00/01918
PRIOR TILING DATE: 2000-03-28
PRIOR APPLICATION UNMBER: JP 11-371382
PRIOR FILING DATE: 1999-12-27
PRIOR APPLICATION UMBER: JP 11-371382
PRIOR FILING DATE: 1999-06-24
PRIOR APPLICATION UMBER: JP 11-178999
PRIOR FILING DATE: 1999-06-24
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; ORGANISM: Homo sapiens
US-10-104-047-3729
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                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                y Match 18.2%; Score 198; DB 15; Length 127;
Local Similarity 36.8%; Pred. No. 3.7e-14;
hes 46; Conservative 19; Mismatches 54; Indels
122 LGEIL 126
                                                        176 LRRIL 180
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                                                                                                                                                                      120 KAEPEGLRRI--HRAG--VQIAIMTFKDYFYCWNTFVENHERTFKAWEGIHENSVRISRQ 175
                                                                                                                 63 DYQ-EALRSICQKRDGPRATMKIMNYDEFQHCWSKFVYSQRELFEPWNNLPKYYILLHIM 121
                                                                                                                                                                                                                                                                                       61 FLRYISDW-DLDPGRCYRVTWFTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYFCEDR 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81 FTSWSPCYDCARHVADFLRGNPNLSLRIFTARLYFCEDRKAEPEGLRRLHRAGVQIAIMT 140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24 RRETYLCYVVK--RRDSATSFSLDFGYLRNKNGCHVELLFL-RYISDWDLDPGRCYRVTW 80
                                                                                                                                                                                                                               3 FFHWFSKWRKLHRDQBYEVTWYISWSPCTKCTRDMATFLAEDPKVTLTIFVARLYYFWDP 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FISWSPCGECSRAITEFLSRHPYVTLFIYIARLYHHTDQR-NRQGLRDLISSGVTIQIMT 145
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Search completed: March 18, 2004, 06:09:26
Job time : 57 secs